<b>ስ</b> ታስ	980	gan	ተሰበሰ	1010	1020
AGETHITIACIO	. まりまいでででなる . まりまいでででする	シモベ - エコンシャルムン APPC	יייריבייורערטבר	GGGTAGCAGG	
				CCCATCGTCC	
1030	1040	1050	1060	1070	1080
				TTKAKAKKA	
Cigcinccan	TCAUCAAAGT	CCCCCTTCTT	TTAGTCCTTT	TTTTTTTAA	AATCTTCGTA
,			****		
		1110			1140
				TTCTCCCCCA	
AAGTTETTEG	TICTACCITA	TARACATOTT	TIGIÇÇAÇGA	AAGAGGGGT	GGTACGCTGG
1150	1160	1170	1180	1190	1200
				CATTCAAAAC	
				GTAAGTTTTG	
accerrana	TOMETHINE	TOICITALOG	SKKST A1 PAUT	ATTENNE TETA	1919191919
1210	1220	1230	1240	1250	1260
ACACACACAC.	ACACACACAC	ACACACACAC	ACACACACAT	GTTTTCTTCC	CTCCCTCCAC
TGTGTGTGTG	TGTGTGTGTG	THTOTOTOTO	TGTGTGTGTA	CAAAAGAAGG	GAGGGAGGTG
1000					
1270			1300		1320
				CACTOTAGAA	
ANGUAGGELA	AGAGACACCA	GGGTTTCTCT	ACIGUTATAA	CTGACATCTT	TAGITATAGI
3330	1340	1350	1360	1370	1380
	CATTOTICACAC		PUNCSINALISIA.	TINTCATTAA	GGTTTTCAATT
ביבורות שונים ביבורים	CIUS CENCOCIO	CONTITIONS	TELESCENCE AND A SOME	ARTAGTAATT	KAUPDAKADD
G**** ** 7000	ATMINUTE TO				
1390		1410	1420	1430	1440
CTTGCCACGT	GTGGGTTTTA	ACCTTTTTAG	GGATTTTTAT	CTAGCGGCAC	TCACCTGCTT
GAACGGTGCA	CACCCAAAAT	TCCAAAAATC	CCTAVAVATA	. GALICGCCGIG	<b>AGTGGACGAA</b>
* * * * * *			4400	1490	1500
1450		1470	1480		
CCCTGTGAAT	GITTAGARTI	CACTGGGCTT	GGTCAGCTAA	TGGAAATGAT	CIMIGGITIG
GUGALACTIA	CMAGTCTTAA	GTGACUCGAA	CCAGICGATT	ACCTTTACTA	GATALCAAAC
1510	1520	1530	1540	1550	1560
				GGAGGGAGAA	
TGAATTTACA	العلىالمالك تكاملتان		הריוויוזייוערוייוערונייזיני הריווייזייועריייזיינערונייזיני	CCTCCCTCTT	TCTCCCCTTC
· 1570	1580	1590	1600	1610	1620
GGAAAACTGC	CITTIATECC	TATTGCTACI	CTAACATIT	GICTCTCACC	TTCCACTTGG
CCTTTTGACC	GAAAATACGG	ATAACGATGA	GATTGTAAA	A CAGAGAGTGG	AAGGTGAACC
			. 920	1.670	0831
TACTUMENT	GAMAGACTIC	ALAGAAAGCT	COUNTRACTOR .	CAGGGATAGG	
WIGNIGHTAG	CITICICAL	: TAICLITCG	f CCCTCGGTC	a mancemance	TOUTOROROR
1690	1700	1710	172	1730	1740
					AGCCTGCTAG
CACACACAC	CCCCCCCAC	י כפיורמייזירמי	CICCGAATC	r CIGICICITY	TOGGACGATO
		. UWLLOVAKY.			
1750			178		
AGAYCATGA	F CTTYCTTTG	A GACCCCTAG	r gctnacaga	A ATAGTICCT	A ACCAGGINGC
TOTROTACTO	GAARGAAAC	r cregegard	A CGATTGTCC	T TATCAAGGA	TGGTCCATCG
1810			0 194		0 . 1860
TGTGGTCAC	TEACTCECC	r ggaagscom	G GCTTTGTCT	T TITGUTTGC	T GTGCAGCCTT
ACACCAGIG	C ACTUNGCCG	a ccttcsgga	c cgaaacaga	a aaacgaacg	a cacgroggaa

## FIG. 1

•	6.7					-
. 1870	1880	1890	1900	1910	1920	
Gaacaaacac	CCTGGCCTCT	TIGAACCCCA	CTATTTCTCA	CCCCTCAGAT	CARGAGGES	
				CGGGAGTCTA		
6110111010	COUNTRIBUTE.	WWCT.TREGG.	MAINIME T	COCCHUTCTY	CITCITICALL	
1930	9848	4050	2000	1020	4000	
	1940	1950	1960		1980	
1031300130	GALGATACIG	APGGGTTCAA	GIGAACTAGG	GCAGAGGGTG	GANGGITTIG	
AUCATIGGAAC	CTCCTATGAC	TACCCAAGTT	CACTTGATCC	CGTCTCCCAC	CTTCCAAAAC	
,						
1990					2040	•
TAACCATAAA	CIGARCIGGG	GTGTTGGTTA	GINAGTAGCC	ATGAA!!ACCA	TAAAAATATATC	
ATTGGTATTT	CACTTCACCC	CACAACCAAT	CATTICATICGG	TACTTATGGT	ATTTTTATAG	
2050					2100	
TGTCAGGTGG	CCAGAGCATU	ACTORGUECA	GAACACAACG	GCCCACTCAG	AACAEGEGGA	
ACAGTCCACC	GGTCTCGTAG	TGACACAAGT	CITCICITOC	CUCGTGAGIC	TTGTGCGCCT	
2110	***			·2150	2160	
Caattcaaag	GCACCAACCT	COGTGCTTCC	TACCCGTTGT	TTTGTTACCG	TGTAAACGCA	
GITAACTITC	CCTCCTTCCA	GGCACGNAGG	ATGGGCAACA	AAACAATGGC	ACATTTGCGT	
2170						
ACTYLARCTCT	CGGCACTGAA	CAGGCTTTTG	CIGCAGACCT	GGGGTCTGGA	GGTGPTGTCT	
TCACTTCAGA	GCCGTGACTT	<b>QTCCGAAAAC</b>	GACCTCTGGA	CCCCAGACCT	CCACAACAGA	
2230	2240	2250	2260	2270	2280	
.CYGAGACACA	AAAACTCATC	TTOTTACTAT	<b>GCATAGTAG</b>	TAACCACGGA	GCTCTGAGAT	
GACTUTGTCC	TTTTGAGTAG	AACAATGATA	CCGTATUATO	ATTIGGTGCCT	CCAGACTCTA	
			***************************************			
2290	2300	2310	2320	2330	2340	
<b>AGCCCTGAGC</b>	TGGTGCCGTT	TAGAAAAGTT	TGATGCTTTA	GAAAGAAATC	GTGGCTTAAA	
TUGGGAUTCG	ACCACGGCNA	ATCTTTTCAA	ACTACGAAAT	CTTTCTTTAG	CACCGAATTT	
			,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,		***************************************	
2350	2360	. 2370	2380	2390	2400	
AGAAGCCTAC	CTGGCATGGG	GGCCCATCCT	CTCCAGCCAT	CCGAATCTCA	ATCTGGTCGT	
TCTTCCGATC	GACCOTACCC	CCGGGTAGGA	GAGGTCGGTA	CCCTTAGAGT	TAGACCAGCA	
2410	2420	2430	2440	2450	2460	
GTGCCTAAGA	ATAGAATCCT			GCTTTTTCTT		
CACGCATTCT	TATCTTAGGA	GCCTTACCAT	MADACAGAA	CGAAAAAGAA	GACCCGAACG	
				-		
2470	2480	2490	. 2500	2510	2520	
TGAGGAAGTC	CCAGGCAGCG	TAGACGTCTT	GGGGGTINGGT	CIYGGAAAAA	TCTCCCAAGA	
ACTOCTTCAC	GGTCCGTCGC	ATCTCCAUAA	CCCCCATCCA	GACCCTTTTT	AGAGGGTTCT	•
					•	
. 2530			2560		2580	-
TTTTAGGAGG	CCACCCCCC	GGATCAGANA	CTTUCAGATT	CGGTAGATCG	CTGTAGAGCA	
AAAAT CCTCC	: ccandeecc	CCUACTCTT	GANCCICTAN	GCCATCTAGC	GACATOTOGT	brain 5'-race product).
• 700			Pulnuve tra	necubilonal start s	the (2, - eng of tur	brain 5'-race product).
2590				2630		
ACTCAGACAG	TCGGCGGCCT	GRAGAGGACT	TGTGCAÁACA	CITCCTCTCT	GGACAAGGAG	
TGAGTCTGTC	AGCCGCCGGA	CTICTCCIGA	ACACGITTGI	GAAGGAGAGA	CCIGIFCCIC	
					•	•
2650			2680			
Gaatgcagga	GECCACCEC	TGCAGTACAT	CITGGAGTGI	TGUAGGGATG	TECCTECACT	
CTTACGTCCI	· cccarecces	ACGTCATGTA	GAACCTCACA	ACCTOCCTAC	ACGGACGTGA	
•				nal start site in re		
2710		<b>Z</b> 2730	2740	2750	2760	<b>–</b>
TGTGAAAGCC	CGCCACARGO	ACCACIOCCC	AACCAAGCCC	GGCAGTGCCC	AGTAGATGCA	,
ACACTTTCCC	GCGGTCTTCC	TGCTCCGGGG	TIGGTICGGG	CCGTCACGGG	TCATCTACCT	
		<del> </del>				
2770	2780	2790	2800	2810	2820	
UNUAGCOTCO	CTUCCCCCGG	CGCACACTWO	GCTCCCTGC	COCCCAGGGG	CCTGAGTCTC	
CTCTCGCAGO	GACGGGGCCC	GCGTGTCAW	CCGAGGGACG	CEGGGIECCE	GGACTCAGAG	
			, <del></del>			

## FIG. 1 cont.

Putative translational start site in murine GLP-2 Receptor gene.

	/	•			
		วลรก			
TCCACKCCCA	CGGGATGCGT	CGGCTCTGGG	GCCCTGGGAC	SCCCTTCCTC	TCCCTCCTTC
		GCCGAGACCC			
•					
2890	2900	2910	2920	2930	2940
TGCT'GGTTTC	CATCAACCAA	GTANGAACAG	STITTTTATTC	CTCATTCGTC	TIGITAATAT
ACGACCAAAG.	GTAGTTCGTT	CATTCTTGTC	DIATAAAAT	GAGTAAGCAG	AACAATTATA
2950	2960	2970	2980	2990	3000
TATCAGTTGT	SCATGTTTTC	TGAGTGTAGA	AGCANTITAG	GCCCCGTGTA	GGCAATTIGG
		ACTCACATOR			
		•			
3010	3020	3030	3040	3050	3060
GTAAGAATAA	AACCATATTA	<b>ACRARATIONG</b>	GCTCAACCAC	AACCCCAGTA	GCATTCTGCT
CATTCTTNT	TIGGIATAAT	TCTTTTACTC	CGAGTTGGTG	TIGGGGT'CN'I	CGTAAGACGA
		3090		3110	
		AAAATTTTTAAAA			
Gygninagya	TAAAACCGAC	<b>ተታ</b> ሞኮለልዶልል፻	AKDADAGTTT	AAGACACGTA	ATMAAATGTG
3130	3140	3150	3160	3170	3180
AGCCGAAATT					
TCGGCTTAA	ومتناهماتين				

<sup>3&#</sup>x27;-End of murine GLP-2 Receptor gene sequenced to date.

FIG. 1 cont.

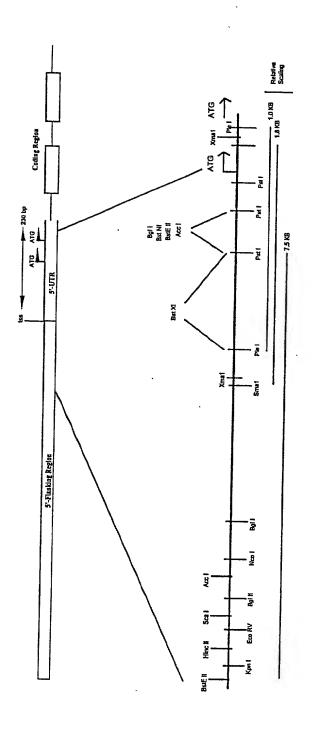
Sequence alignment of the 5' end of the mGLP-2 receptor gene with the 5' end of the cDNA encoding the rat GLP-2R.

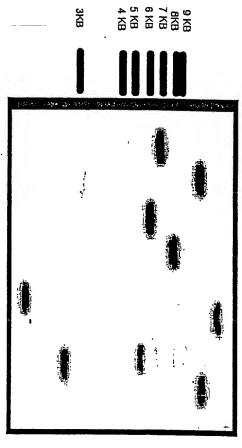
Putative	transcriptional start site.	
*	5'-UTH nearly identical between	
MOUSE GLP-2R RAT GLP-2R	10 20 30 40 50  1 AACACTTCCT CTCTGGACAA GGAGGAGTGC AGGAGGCCAC CGCCTGCAGT  AACACTTCCT CTCTGGACAA GGAGGAGTGC AGGAGGCCAC CGCCTGCAGT	
MOUSE GLP-2R RAT GLP-2R	51 ACATCITEGA ETGITEGAGG GATGTCCCTG CACTTGTGAA AGGGCCCCAG 51 ACATCITEGA GTGTTGGAGG GATGTCCCTG CACTTGTGAA CGGCCCCCAG	
Mouse Glp-2r Rat Glp-2r	101 NAGRACGAGG CCCCAACCAA GCCCGGCAGT GCCCAGTAGA TGCAGAGAGC 101 GACAATGAGG CCCCAACCAA GCCCGGCAGT GCCCAGTAGA TGCAGAGAGC	
MOUSE GLP-2R RAT GLP-2R	151 GTCCT-TGCC CCGAGTGAGG GCACAGCCAG TGGGGCTCCC TGGGGCCCAG 151 CACCTGTGCC CCGAGTGAGG GCACAGCCAG TGGGCATCCC TGAGGCCCAG	
MOUSE GLP 2R MAT GLP-2R	201 GGGCCTGAGT CTCTCCACKC CCAACAGATG CGTCGGCTCT GGGGCCCTGG	
Mouse Glp-2r Rat Glp-2r	251 GACGECCTTC CTCGCCCTGC TTCTGCTGGT TTCCATCAAG CAAGTAAGAA 251 GAGGCCCTTC CTCGCCCTGC TTCTGCTGGT TTCCATCAAG CAAGTTACAG	
MOUSE GLP-2R RAT CLP-2R	310 320 330 340 350 301 CAGATTTTTA TTCCTCATTC GTCTTGTTAA TATTATCAGT TGTGCATGTT 301 GATCGCTCCT CAACGAGACA ACTCAGAAGT COGCTAATTA TAAGGAGAAG	
Upstream initiator ATG codon.		
	Downstream initiator ATG codon.	

Bequence alignment of the 5' end of the mGLP-2 receptor gene with the 5' end of the cDNA encoding the rat GLP-2R.

The 5' end of the cDNA encoding the rat GLP-2R (cloned by 5'-RACE) is presented in alignment with the corresponding region of sequence encoding the murine GLP-2R. The upstream initiator ATG codon is present in the rat sequence, and the downstream initiator ATG codon is conserved between in both the rat and murine sequences encoding the GLP-2R. The sequence corresponding to the putative 5'-UTR (untranslated region) is nearly identical between the rat and murine sequences presented.

## FIG. 2





Hind III

Bam H1

Eco R1

Pvu II

PST I

Sal I

Xba I

Xho I

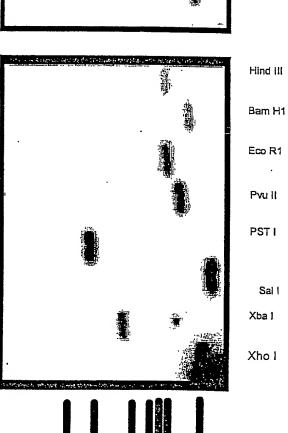


FIG. 5

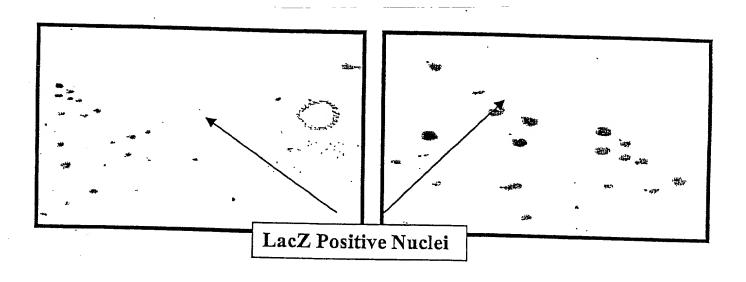


FIG. 6

17

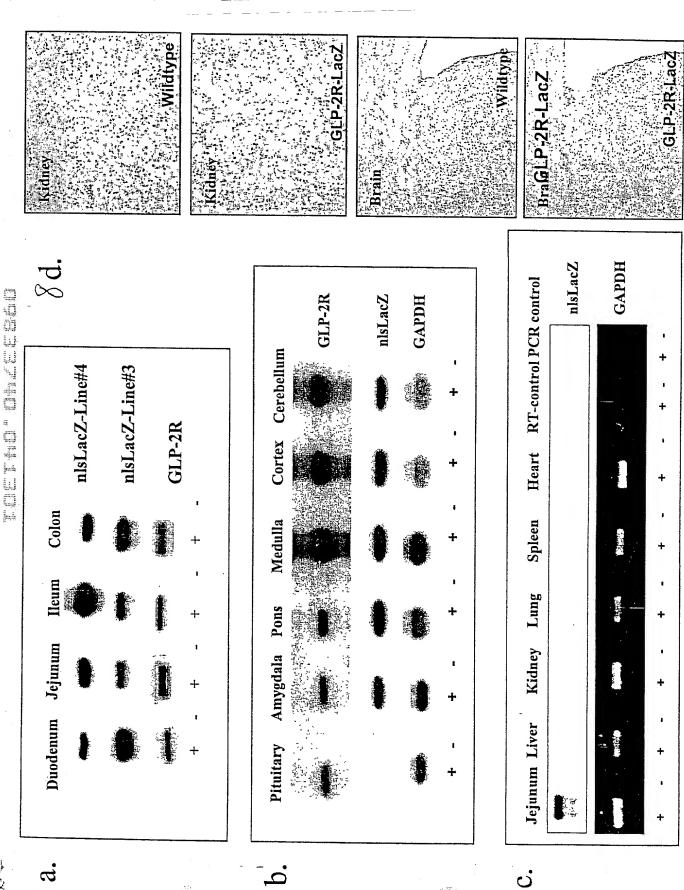
146.00

CCC CGA GTG AGG GCA CAG CCA > × Д, GCA CCC GTG P. R E A gaga ATG AGG CCC CAA CCA AGC CCG GCA GTG CCC AGT AGA TGC AGA GAG 5'-UTR GGC ATC CCT GAG GCC CAG GGG CCC GTT CCT CTC CAC TCC CAA CAG ATG ¥ 0 ပ 0 × S S G P V P L H <u>.</u> > < ۵, S 4 o 0 E A G I P ♦ 5'-end Σ rat GLP-2R cDNA GLP-2R CDNA CDNA rat GLP-2R

--atttttat tcctcattc TCC AIC AAG CAA gtaagagcagttca ttattattat tattatcag gaaagcacag ctgacttagg ga-aggtctg ggaaaaatct occaagattt aggaggga:--ggcggggggatgaggagct tggagattcg gtagatcgct gt---agagc aactcagaca actecgtaga ttgetetaga  $c_{\mathrm{TG}}$ aacactt cctctfgga caaggaggag tgcaggaggc ccgcctcaga cactctcggc gcagcgtgga gaggatttgt gcaaacattt cctctgtgga ccaagaggaa tgcaagagga SSS ည္ဟ ACA ည္ပ 9 ű GGC GCC Ü gtacatott ggagtgttgg agggatgtgo otgcacttgt gaacgggege caggaga ATG AGG caccgcc tgca gtacatott ggagtgttgg agggatgtgc ctgcacttgt gaaagggcgc cagaagg ACG AGG gtgcatctt ggacggctag agagatgtac ccctacttgt gaaggtgcac gaggaag ATG AAG  $\mathbf{c}_{\mathrm{LG}}$ TGG CGC TGG GCA CAG TGG 3 **9**99 GAG CIC CTG CTG CIC TCC AIC AAG CAA gtaagaacag---ggtaggtetg caaggaggaa GTG AGG GGA TCG AGC AGG GCA GGG CCT GGG AGA GGA AGC GCG GGA CTC CTG CCT GGC GTC CAC CGG --- -GT CCC TGC CCC TCT TCC CAA CAG ATG CGT GGG ATG CGT TTC CAC AGG AAG TGC CAA CCA AGC CCG GCA GTG CCC AGT AGA TGC AGA GAG GCA CCC GTG CCC CGA atgtettget ttttettetg ggettgetga ggaagteeca ggeagegtag aegtettggg gcaaacactt cctctctgga TCC ATC AAG CAA KCC CAC CAA CCA AGC CCG GCA GIG CCC AGI AGA IGC AGA GAG C-- ---TCC CIC CAC CIC CAC GTT GII CIG GIL cogcottgit ctitotooto agoctgicaa ggaagtooca NF-Kappa B ----tgaa gaggacttgt TCT CCT GIT CCI CTG CTG CIG CIG GAG AGT CIG TGG GGG ACC වූපු පදුපු GCG GCC CAG GGG CCT CTG CTT CIG CII н CAG 900 TCC ပ္ပ ပ္ပ gteggeggee ------GAG ပ္ပပ္ပ TIC TIC CCT CCT ပ္ပ ပ္ပ GCT GGG AGG CCC ggctgcc tgcg caccgcc tgca CIC AIC GGC ATC GCT GGG AGG GCT GGG ACG ည္ဟ ධ්රී Ü GIG ATG GTW 38 180 -123 mouse GLP-2R human GLP-2R human GLP-2R mouse GLP-2R mouse GLP-2R human GLP-2R mouse GLP-2R human GLP-2R mouse GLP-2R human GLP-2R mouse GLP-2R mouse GLP-2R human GLP-2R human GLP-2R rat GLP-2R rat GLP-2R rat GLP-2R rat GLP-2R cat GLP-2R

ŝ Xho I က nls -LacZ Pstl 15-Kb-GLP-2R Promoter Sma / Sac II in

70



19 60

(%)

ပ တ

